



**S2 Fig. Evidence for recombination within Clade II throughout the *Ophidiomyces ophidiicola* genome.** Log-transformed Phi test p-values calculated using PhiPack with a window size of 50,000 bases and a step size of 25,000 bases (1). Black horizontal dashed line indicates a p-value of 0.05 with Bonferroni correction for 749 different tests. Red vertical dashed lines indicate scaffold boundaries within the reference assembly. Data underlying this figure can be found in OSF: <https://osf.io/fmbh5/>.